OIPE

RAW SEQUENCE LISTING

DATE: 04/05/2001

PATENT APPLICATION: US/09/813,492

TIME: 12:15:06

Input Set : A:\SEQUENCE LISTING.txt Output Set: N:\CRF3\04052001\I813492.raw

ENTERED

```
3 <110> APPLICANT: Labow, Mark A.
      4
              Mickanin, Craig Stephen
              Bhatia, Umesh
      7 <120> TITEE OF INVENTION: MAMMARY GLAND CHEMOKINE
     10 -: 130> FILE REFERENCE: 12345
C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/813,492
C--> 12 <141> CURRENT FILING DATE: 2001-03-21
     12 <160> NUMBER OF SEQ ID NOS: 2
     14 <170> SOFTWARE: FastSEQ for Windows Version 4.0
     16 <210> SEQ ID NO: 1
     17 <211> LENGTH: 2017
     18 -: 212> TYPE: DNA
     14 <213 · ORGANISM: HUMAN
     21 ₹400> SEQUENCE: 1
     22 tagataccet gaacacetee cagggegggg ceaectgget taettiteet etgeaettte
                                                                                60
     23 totgtgooda aggadadett tageoteatt tootgatoga adagootead tigtgitigot
                                                                               120
     24 gteagtgeea gtagggeagg caggaatgea geagagagga etegeeateg tggeettgge
     25 tytotytycy geoctacaty coteagaago cataottoco attycotoca gotyttycac
                                                                               240
     26 ggaggtttca catcatattt ccagaagget cetggaaaga gtgaatatgt gtcgcatcca
                                                                                300
                                                                                360
     27 gagagetgat ggggattgtg aettggetge tgteateett eatgteaage geagaagaat
                                                                               420
     28 otgtgtoago oogoacaaco atactgttaa goagtggatg aaagtgcaag otgocaagaa
     29 haatggtaaa ggaaatgttt gocacaggaa gaaacaccat ggcaagagga acagtaacag
     30 ggcacatcag gggaaacacg aaacatacgg ccataaaact ccttattaga gagtctacag
                                                                               540
     31 ataaatotao agagacaatt ootoaagtgg acttggcoat gattggttgt aagtttatoa
                                                                               600
     32 totgaattot oottattgta gacaacagaa caaaacaaaa tattggtttt taaaaaaatga
                                                                               660
     33 acaattytyo yytätyoaaa tytäyeeaat aatataotea aaoteetyyy eteääyeyät
                                                                               720
     34 ectoccaect tagectocca aagtactggg attataggty tgagccaeag tgeetggeet
     35 aattattito tigigatoaa attoaggitti aatgittitig gitaagaatt tootaogiga
                                                                               840
                                                                               900
     36 attogtique ttattttique atttaqaqtt cataaatatt agggtttatt ttotaaataq
     37 aatagtitaa actaaatata actteaaaac gtetagtitg agtagetaec gtigtitgga
                                                                               460
     38 ttgaadtttt otgataotga aaagaacaaa aagootgoot ttotgoocag aacottttgo
                                                                              1020
     39 cteccceagt cagttettgg ageageacta gttaggggee cagagttegg cettetgtgt
                                                                               1080
     40 ggtgatttta egetetgeet aaacaaggag ootacatett ttageteeta ttecaceett
                                                                              1140
     41 eteacacett tttgttgttg tttggttgtt ttttttttgag acagagtote actetgttge
                                                                              1200
                                                                              1260
     42 ocaggotiga gigologigo acaatotogg etcatigoaa ecicogeeto eegegiteaa
     43 gigattotet igeeicagee teeeaagiaa eigatattae aggegeeeag eeaeeacaee
                                                                              1320
     44 cogetquatt ttgtattttt agtaqaqacq qqqttttccc acqttqqccq qqctqqtctc
     45 aaaetettga eeteaagtga accaeeegee tytyeeteee aaagtgetyg aattaeeage
     46 gigagecace algoeggget cacaegitty agittgatace altigipecat tectofitty
                                                                              1500
                                                                              1560
     47 geotettitt tigteeataga ggotteaaga tagataggta agageeeagt agtigteeata
     48 agaageeaat agagageagg ageeaettta teaggtggea ggtgteeegg geeteeetge
                                                                              1620
     49 tggotagtoc caageggtgg tgttgccagg atgtottgga ggtgataatg ggacacacag
                                                                              1680
     50 aggeactgay tetecatagy ttaaaatgee accaaaactg geetttgeet aatateeete
                                                                              1740
     51 attgaetatt tageatttaa titatitatt tieetgaeat tietgeaage titigtattta
                                                                              1800
     52 tattteeaet ttatagatga ggaaatttga ggetettaga ggtaaaatga ettgeeeagg
                                                                              1860
     53 tcacacagga agtggcagag acaagctttt taaataagaa aaaattaata aaatataata
                                                                              1920
     54 tgagagtaac ttaaaatatt aataaaccac aattttaaat taattaaccg tgataaccaa
                                                                              1480
```

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55	is cattaataaa agttaagata ccaaaaaaaa aaaaaaa														2	2017	
57	-:210	)> SI	EQ II	ON C	2												
58	-:21:	L: LE	ENGTI	H: 13	27												
59	59 -: C12:- TYPE: PRT																
60	50 -: 213: ORGANISM: HUMAN																
62	62 - (400) - SEQUENCE: 2																
63	Met	Gln	Gln	Arg	Gly	Leu	Ala	Ile	Val	Ala	Leu	Ala	Val	Cys	Ala	Ala	
64	1				5					10					15		
65	Leu	His	Ala	Ser	Glu	Ala	Ile	Leu	Pro	Ile	Ala	Ser	Ser	Cys	Cys	Thr	
66		•		20					25					30			
67	Glu	Val	Ser	His	His	Ile	Ser	Arg	Arg	Leu	Leu	Glu	Arg	Val	Asn	Met	
68			35					40					45				
69	Cys	Arg	Ile	Gln	Arg	Ala	Asp	Gly	Asp	Cys	Asp	Leu	Ala	Ala	Val	Ile	
70		50					55					60					
71	Leu	His	Val	Lys	Arg	Arg	Arg	Ile	Cys	Val	Ser	Pro	His	Asn	His	Thr	
72	65					70					75					80	
73	Val	Lys	Gln	Trp	Met	Lys	Val	Gln	Ala	Ala	Lys	Lys	Asn	Gly	Lys	Gly	
74					85					90					95		
75	Asn	Val	Cys	His	Arg	Lys	Lys	His	His	Gly	Lys	Arg	Asn	Ser	Asn	Arg	
76				100					105					110			
77	Ala	His	Glņ	Gly	Lys	His	Glu	Thr	Tyr	Gly	His	Lys	Thr	Pro	Tyr		
78			115					120					125				



DATE: 04/05/2001 TIME: 12:15:07

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L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date